



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

10/517,214

Source:

PCT

Date Processed by STIC:

12-22-04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER
VERSION 4.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):
U.S. Patent and Trademark Office, 220 20th Street S., Customer Window, Mail Stop Sequence, Crystal Plaza Two, Lobby, Room 1B03, Arlington, VA 22202

Revised 05/17/04



PCT

RAW SEQUENCE LISTING

DATE: 12/22/2004

PATENT APPLICATION: US/10/517,214

TIME: 09:16:36

Input Set : A:\3065 USOP seq.txt

Output Set: N:\CRF4\12222004\J517214.raw

2 <110> APPLICANT: MAEKAWA, Tsuyoshi
 3 HARA, Ryoma
 4 ODAKA, Hiroyuki
 5 MIZUFUNE, Hideya
 6 FUKATSU, Kohji
 8 <120> TITLE OF INVENTION: 1,2-Azole Derivatives with Hypoglycemic and Hypolipidemic Activity
 10 <130> FILE REFERENCE: 3065 USOP
 C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/517,214
 C--> 12 <141> CURRENT FILING DATE: 2004-11-23
 12 <150> PRIOR APPLICATION NUMBER: PCT/JP03/06389
 13 <151> PRIOR FILING DATE: 2003-05-22
 15 <150> PRIOR APPLICATION NUMBER: JP 2002-151405
 16 <151> PRIOR FILING DATE: 2002-05-24
 18 <150> PRIOR APPLICATION NUMBER: JP 2002-287161
 19 <151> PRIOR FILING DATE: 2002-09-30
 21 <150> PRIOR APPLICATION NUMBER: JP 2003-016748
 22 <151> PRIOR FILING DATE: 2003-01-24
 24 <160> NUMBER OF SEQ ID NOS: 12
 26 <210> SEQ ID NO: 1
 27 <211> LENGTH: 34
 28 <212> TYPE: DNA
 29 <213> ORGANISM: Artificial Sequence
 W--> 30 <220> FEATURE:
 31 <223> OTHER INFORMATION: Primer/probe
 33 <400> SEQUENCE: 1
 34 aacggtacct cagccatgga gcagcctcag gagg 34
 36 <210> SEQ ID NO: 2
 37 <211> LENGTH: 34
 38 <212> TYPE: DNA
 39 <213> ORGANISM: Artificial Sequence
 W--> 40 <220> FEATURE:
 41 <223> OTHER INFORMATION: Primer/probe
 43 <400> SEQUENCE: 2
 44 taagtcgacc cgtagtagta tgctccttgta gatc 34
 46 <210> SEQ ID NO: 3
 47 <211> LENGTH: 33
 48 <212> TYPE: DNA
 49 <213> ORGANISM: Artificial Sequence
 W--> 50 <220> FEATURE:
 51 <223> OTHER INFORMATION: Primer/probe
 53 <400> SEQUENCE: 3
 54 ttagaattcg acatggacac caaacatttc ctg 33
 56 <210> SEQ ID NO: 4

Does Not Comply
Corrected Diskette Needed

(pg. 2)

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/517,214

DATE: 12/22/2004

TIME: 09:16:36

Input Set : A:\3065 USOP seq.txt

Output Set: N:\CRF4\12222004\J517214.raw

57 <211> LENGTH: 33
58 <212> TYPE: DNA
59 <213> ORGANISM: Artificial Sequence
W--> 60 <220> FEATURE:
61 <223> OTHER INFORMATION: Primer/probe
63 <400> SEQUENCE: 4
64 cccctcgagc taagtcattt ggtgcggcgc ctc 33
66 <210> SEQ ID NO: 5
67 <211> LENGTH: 36
68 <212> TYPE: DNA
69 <213> ORGANISM: Artificial Sequence
W--> 70 <220> FEATURE:
71 <223> OTHER INFORMATION: Primer/probe
73 <400> SEQUENCE: 5
74 tcgacagggg accaggacaa aggtcacgtt cgggag 36
76 <210> SEQ ID NO: 6
77 <211> LENGTH: 36
78 <212> TYPE: DNA
79 <213> ORGANISM: Artificial Sequence
81 <220> FEATURE:
82 <223> OTHER INFORMATION:
W--> 84 <400> 6
85 tcgactcccg aacgtgacct ttgtcctggt cccctg 36
87 <210> SEQ ID NO: 7
88 <211> LENGTH: 28
89 <212> TYPE: DNA
90 <213> ORGANISM: Artificial Sequence
W--> 91 <220> FEATURE:
92 <223> OTHER INFORMATION: Primer/probe
94 <400> SEQUENCE: 7
95 cccagatctc cccagcgtct tgtcattg 28
97 <210> SEQ ID NO: 8
98 <211> LENGTH: 28
99 <212> TYPE: DNA
100 <213> ORGANISM: Artificial Sequence
W--> 101 <220> FEATURE:
102 <223> OTHER INFORMATION: Primer/probe
104 <400> SEQUENCE: 8
105 tcaccatggt caagctttta agcggggtc 28
107 <210> SEQ ID NO: 9
108 <211> LENGTH: 33
109 <212> TYPE: DNA
110 <213> ORGANISM: Artificial Sequence
W--> 111 <220> FEATURE:
112 <223> OTHER INFORMATION: Primer/probe
114 <400> SEQUENCE: 9
115 gtgggtaccg aaatgacat gggtgacaca gag 33
117 <210> SEQ ID NO: 10
118 <211> LENGTH: 33

pls explain source
of genetic material.
See item # 11
on errn summary
sheet.

RAW SEQUENCE LISTING

DATE: 12/22/2004

PATENT APPLICATION: US/10/517,214

TIME: 09:16:36

Input Set : A:\3065 US0P seq.txt

Output Set: N:\CRF4\12222004\J517214.raw

119 <212> TYPE: DNA
120 <213> ORGANISM: Artificial Sequence
W--> 121 <220> FEATURE:
122 <223> OTHER INFORMATION: Primer/probe
124 <400> SEQUENCE: 10
125 ggggtcgacc aggactctct gctagtacaa gtc 33
127 <210> SEQ ID NO: 11
128 <211> LENGTH: 33
129 <212> TYPE: DNA
130 <213> ORGANISM: Artificial Sequence
W--> 131 <220> FEATURE:
132 <223> OTHER INFORMATION: Primer/probe
134 <400> SEQUENCE: 11
135 cgtcgacccg gcggcccat ggacctgccc ccg 33
137 <210> SEQ ID NO: 12
138 <211> LENGTH: 33
139 <212> TYPE: DNA
140 <213> ORGANISM: Artificial Sequence
W--> 141 <220> FEATURE:
142 <223> OTHER INFORMATION: Primer/probe
144 <400> SEQUENCE: 12
145 catcgattag cagtggcggtt acttctggga ctt 33
W--> 146 ??
W--> 148 ??
W--> 150 ??
W--> 152 ??

*PLS
delete*

VERIFICATION SUMMARY

DATE: 12/22/2004

PATENT APPLICATION: US/10/517,214

TIME: 09:16:37

Input Set : A:\3065 USOP seq.txt

Output Set: N:\CRF4\12222004\J517214.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:30 M:283 W: Missing Blank Line separator, <220> field identifier
L:40 M:283 W: Missing Blank Line separator, <220> field identifier
L:50 M:283 W: Missing Blank Line separator, <220> field identifier
L:60 M:283 W: Missing Blank Line separator, <220> field identifier
L:70 M:283 W: Missing Blank Line separator, <220> field identifier
L:84 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:6,Line#:82
L:91 M:283 W: Missing Blank Line separator, <220> field identifier
L:101 M:283 W: Missing Blank Line separator, <220> field identifier
L:111 M:283 W: Missing Blank Line separator, <220> field identifier
L:121 M:283 W: Missing Blank Line separator, <220> field identifier
L:131 M:283 W: Missing Blank Line separator, <220> field identifier
L:141 M:283 W: Missing Blank Line separator, <220> field identifier
L:146 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:148 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:12
L:148 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:150 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:12
L:150 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:152 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:12
L:152 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1

STATISTICS SUMMARY

PATENT APPLICATION: US/10/517,214

DATE: 12/22/2004

TIME: 09:16:37

Input Set : A:\3065 US0P seq.txt

Output Set: N:\CRF4\12222004\J517214.raw

Application Serial Number: US/10/517,214

Alpha or Numeric or Xml: Numeric

Application Class:

Application File Date: 11-23-2004

Art Unit: PCT

Software Application:

Total Number of Sequences: 12

Total Nucleotides: 394

Total Amino Acids: 0

Number of Errors: 0

Number of Warnings: 19

Number of Corrections: 2

MESSAGE SUMMARY

258 W: 1 (Mandatory Feature missing)

270 C: 1 (Current Application Number differs)

271 C: 1 (Current Filing Date differs)

283 W: 11 (Missing Blank Line separator)

334 W: 4 ((2) Invalid Amino Acid in Coding Region)

336 W: 3 (Invalid Amino Acid Number in Coding Region)